

**WHAT IS CLAIMED IS:**

1. A method for creating a target single-stranded region in double-stranded DNA comprising:

5 (a) nicking at least one site bordering the target region in double-stranded DNA with at least one site-specific nicking endonuclease; and

(b) subjecting the nicked DNA to conditions where the target region is selectively denatured.

10 2. The method of claim 1 wherein the target single-stranded region comprises a gap in the double-stranded DNA and wherein the method comprises:

15 (a) nicking at least two sites bordering the target region in one strand of double-stranded DNA with at least one site-specific nicking endonuclease; and

(b) subjecting the nicked DNA to conditions where the target region is selectively denatured.

20 3. The method of claim 1 wherein the target single-stranded region comprises at least one terminus in the double-stranded DNA and wherein said method comprises:

25 (a) nicking at least one site bordering the target region in a first strand of the double-stranded DNA with at least one site-specific nicking endonuclease, wherein the second strand of the double-stranded DNA has at least one break bordering the target region; and

(b) subjecting the resulting DNA to conditions where the target region is selectively denatured.

4. The method of claim 3 wherein the break in the second strand is pre-existing.

5. The method of claim 3 wherein the break in the second strand is produced by a site-specific endonuclease.

6. A method of joining nucleic acid molecules comprising creating a target single-stranded DNA region in a first molecule in accordance with the method of claim 1 and adding a second molecule containing a nucleic acid single-stranded region complementary to that contained on the first molecule.

7. A method of joining nucleic acid molecules comprising creating a first DNA molecule with a target single-stranded DNA region of at least three nucleotides in accordance with the method of claim 1 and adding a second molecule containing a nucleic acid single-stranded region complementary to the target single-stranded region at the first molecule.

8. A method of joining nucleic acid molecules comprising creating a first DNA molecule with a target single-stranded DNA region of at least nine nucleotides in accordance with the method of claim 1 and adding a second

molecule containing a nucleic acid single-stranded region complementary to the target single-stranded region of the first molecule.

5           9.    A method of joining nucleic acid molecules comprising creating a first DNA molecule with a target single-stranded DNA region of at least 12 nucleotides in accordance with the method of claim 1 and adding a second molecule containing a nucleic acid single-stranded region  
10 complementary to the target single-stranded region of the first molecule.

          10.   A method of joining nucleic acid molecules comprising creating a first DNA molecule with a target single-stranded DNA region of at least 18 nucleotides in accordance  
15 with the method of claim 1 and adding a second molecule containing a nucleic acid single-stranded region complementary to the target single-stranded region of the first molecule.

20           11.   A method of attaching a molecular probe to DNA comprising annealing a complementary single-stranded nucleic acid region on the probe to a single-stranded DNA region produced by the method of claim 1.

25           12.   A method for the purification of a specific DNA fragment comprising annealing a single-stranded DNA region

produced by the method of claim 1 to a complementary single-stranded nucleic acid attached to a solid support.

5           13. A method for producing a branched nucleic acid molecule comprising:

          (a) producing a single-stranded gap in double-stranded DNA according to the method of claim 2; and

          (b) annealing to the single-stranded region of step (a) a second nucleic acid molecule containing a single-stranded  
10           region complementary thereto.

          14. A nucleic acid molecule comprising of at least two fragments joined by single-stranded termini produced by the method of claim 3.

15           15. A nucleic acid molecule which comprises three or more fragments joined by single-stranded termini produced by the method of claim 3.

20           16. A nucleic acid molecule which comprises at least two fragments joined by single-stranded termini, in which at least one terminus is produced by the method of claim 3.

25           17. A circular nucleic acid molecule produced by joining at least two fragments containing at least one terminus produced by the method of claim 3.

18. A DNA vector produced by joining fragments containing at least one terminus produced by the method of claim 3.

5 19. A method for assembling a vector with multiple, interchangeable parts comprising element sets containing:

(a) a replication origin and associated control sequences; and

(b) a selectable marker; and

10 (c) one or more of the following element sets:

(i) promoters and associated control elements;

(ii) coding sequences allowing production of fusion proteins;

(iii) transcription terminators;

15 (iv) regulatory proteins; and

(v) gene coding regions;

wherein each element set has unique common cohesive termini, allowing joining of elements within an ordered assembly, and interchange of elements in each set in the assembly.

20

20. A method for generating a DNA fragment with specific single-stranded termini comprising (a) inserting a target DNA fragment between two sets of site-specific nicking sites, wherein said sites are situated to generate single-stranded termini in accordance with the method of claim 3, and (b) digesting the resulting construct with the cognate site-specific nicking endonuclease or endonucleases to

25

release the target DNA fragment containing the single-stranded termini.

21. A method for generating a DNA fragment with  
5 specific single-stranded termini comprising (a) inserting a  
target DNA fragment between two sets of N.BstNBI sites,  
wherein such sites are situated to generate single-stranded  
termini in accordance with the method of claim 3, and (b)  
digesting the resulting construct with N.BstNBI endonuclease  
10 to release the target DNA fragment containing the added  
single-stranded termini.